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RAW SEQUENCE LISTING

DATE: 06/21/2002

PATENT APPLICATION: US/10/023,775B

TIME: 14:54:54

Input Set : A:\PC10959A Sequence Listing.ST25.txt

Output Set: N:\CRF3\06212002\J023775B.raw

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3 <110> APPLICANT: Pfizer Ltd. (EP(GB) only)
4 Pfizer Inc. (US, JP, EB except GB)
5 Fidock, Mark David
7 <120> TITLE OF INVENTION: Novel Polypeptide
9 <130> FILE REFERENCE: PC10959AGPR
11 <140> CURRENT APPLICATION NUMBER: 10/023,775B
12 <141> CURRENT FILING DATE: 2001-12-18
14 <150> PRIOR APPLICATION NUMBER: GB 0030854.4
15 <151> PRIOR FILING DATE: 2000-12-18
17 <150> PRIOR APPLICATION NUMBER: US 60/260,590
18 <151> PRIOR FILING DATE: 2001-01-09
20 <150> PRIOR APPLICATION NUMBER: US 60/296,660
21 <151> PRIOR FILING DATE: 2001-06-07
23 <150> PRIOR APPLICATION NUMBER: GB 0111031.1
24 <151> PRIOR FILING DATE: 2001-05-04
26 <160> NUMBER OF SEQ ID NOS: 4
28 <170> SOFTWARE: PatentIn version 3.1
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31 <211> LENGTH: 1014
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo sapiens
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42 aaaatgagac cttggaagag cagcaccatc attatgctga acctggcctg cacagatctg 240
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54 tcagcctgtc tcgacctcac cagttcggat gaactcaata ctattaagtg gtacaacctc 600
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60 aggctaacca ttctgtact ctttgcatth tacgtatggt ttttaccctt ccatactttg 780
62 agggctcatc ggatcgaaac tcgctgtgct tcaatcagtt gttccattga gaatcagatc 840
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66 ctatatgtgg tggtcagcga caactttcag caggctgtct gctcaacagt gagatgcaaa 960
68 gtaagcggga accttgagca agcaaagaaa attagttact caaacaaccc ttga 1014
71 <210> SEQ ID NO: 2
72 <211> LENGTH: 337
73 <212> TYPE: PRT
74 <213> ORGANISM: Homo sapiens

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76 <400> SEQUENCE: 2

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83          20          25          30
86 Met His Tyr Leu Pro Val Ile Tyr Gly Ile Ile Phe Leu Val Gly Phe
87          35          40          45
90 Pro Gly Asn Ala Val Val Ile Ser Thr Tyr Ile Phe Lys Met Arg Pro
91          50          55          60
94 Trp Lys Ser Ser Thr Ile Ile Met Leu Asn Leu Ala Cys Thr Asp Leu
95 65          70          75          80
98 Leu Tyr Leu Thr Ser Leu Pro Phe Leu Ile His Tyr Tyr Ala Ser Gly
99          85          90          95
102 Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Ser
103          100          105          110
106 Phe His Phe Asn Leu Tyr Ser Ser Ile Leu Phe Leu Thr Cys Phe Ser
107          115          120          125
110 Ile Phe Arg Tyr Cys Val Ile Ile His Pro Met Ser Cys Phe Ser Ile
111          130          135          140
114 His Lys Thr Arg Cys Ala Val Val Ala Cys Ala Val Val Trp Ile Ile
115 145          150          155          160
118 Ser Leu Val Ala Val Ile Pro Met Thr Phe Leu Ile Thr Ser Thr Asn
119          165          170          175
122 Arg Thr Asn Arg Ser Ala Cys Leu Asp Leu Thr Ser Ser Asp Glu Leu
123          180          185          190
126 Asn Thr Ile Lys Trp Tyr Asn Leu Ile Leu Thr Ala Thr Thr Phe Cys
127          195          200          205
130 Leu Pro Leu Val Ile Val Thr Leu Cys Tyr Thr Thr Ile Ile His Thr
131          210          215          220
134 Leu Thr His Gly Leu Gln Thr Asp Ser Cys Leu Lys Gln Lys Ala Arg
135 225          230          235          240
138 Arg Leu Thr Ile Leu Leu Leu Leu Ala Phe Tyr Val Cys Phe Leu Pro
139          245          250          255
142 Phe His Ile Leu Arg Val Ile Arg Ile Glu Ser Arg Leu Leu Ser Ile
143          260          265          270
146 Ser Cys Ser Ile Glu Asn Gln Ile His Glu Ala Tyr Ile Val Ser Arg
147          275          280          285
150 Pro Leu Ala Ala Leu Asn Thr Phe Gly Asn Leu Leu Leu Tyr Val Val
151          290          295          300
154 Val Ser Asp Asn Phe Gln Gln Ala Val Cys Ser Thr Val Arg Cys Lys
155 305          310          315          320
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162 Pro

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167 <211> LENGTH: 33

168 <212> TYPE: DNA

169 <213> ORGANISM: Homo sapiens

171 <400> SEQUENCE: 3

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176 <211> LENGTH: 30
177 <212> TYPE: DNA
178 <213> ORGANISM: Homo sapiens
180 <400> SEQUENCE: 4
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VERIFICATION SUMMARY

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